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OM protein - protein search, using sw model

Run on: February 1, 2005, 14:23:28 ; Search time 147 Seconds
(without alignments)
594.776 Million cell updates/sec

Title: US-10-629-329A-2
Perfect score: 1322
Sequence: 1 MSGCDAGEGDCCRRCAQD.....SMKKVGLDPSQLPVGNGIV 242

Scoring table: BLOSUM62
Gap 10.0 , Gapext 0.5

Searched: 1608061 seqs, 361289386 residues

Total number of hits satisfying chosen parameters: 1608061

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Database : Published Applications AA.*
1: /cgn2_6/ptodata/1/pubpaa/US07_PUBCOMB.pdb.*
2: /cgn2_6/ptodata/1/pubpaa/PCT_NEW_PUB.pdb.*
3: /cgn2_6/ptodata/1/pubpaa/US06_NEW_PUB.pdb.*
4: /cgn2_6/ptodata/1/pubpaa/US06_PUBCOMB.pdb.*
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6: /cgn2_6/ptodata/1/pubpaa/PCTUS_PUBCOMB.pdb.*
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12: /cgn2_6/ptodata/1/pubpaa/US09_NEW_PUB.pdb.*
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18: /cgn2_6/ptodata/1/pubpaa/US11_NEW_PUB.pdb.*
19: /cgn2_6/ptodata/1/pubpaa/US60_NEW_PUB.pdb.*
20: /cgn2_6/ptodata/1/pubpaa/US60_PUBCOMB.pdb.*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Match	Length	ID	Description
1	1322	100.0	242	US-10-629-329A-2	Sequence 2, Appli
2	1307	98.9	242	US-10-220-381-2	Sequence 2, Appli
3	1296	98.0	242	US-10-381-710-4	Sequence 4, Appli
4	1239.5	93.8	241	US-10-381-710-2	Sequence 2, Appli
5	1239.5	93.8	241	US-10-629-329A-4	Sequence 4, Appli
6	573.5	43.4	529	US-10-437-963-195546	Sequence 195546
7	563	42.6	256	US-10-424-599-157170	Sequence 157170
8	563	42.6	517	US-10-425-115-253963	Sequence 253963
9	563	42.6	524	US-10-425-114-64486	Sequence 64486
10	555	42.0	497	US-10-425-114-65135	Sequence 65135
11	553	41.8	517	US-10-767-701-45914	Sequence 45914
12	549	41.5	522	US-10-424-599-273717	Sequence 273717
13	549	41.5	540	US-10-425-114-46271	Sequence 46271

14	514.5	38.9	594	17	US-10-425-115-253964	Sequence 253964
15	467.5	35.4	459	15	US-10-425-114-61505	Sequence 61505
16	415	31.4	533	15	US-10-425-114-57875	Sequence 57875
17	312	23.6	64	14	US-10-029-386-28983	Sequence 28983
18	292	22.1	128	16	US-10-767-701-59226	Sequence 59226
19	228	17.2	59	14	US-10-106-698-5148	Sequence 5148
20	214	16.2	202	14	US-10-156-761-14187	Sequence 14187
21	195	14.8	212	15	US-10-282-122A-45375	Sequence 45375
22	193	14.6	204	15	US-10-282-122A-69836	Sequence 69836
23	182	13.8	204	15	US-10-282-122A-55860	Sequence 55860
24	166	12.6	205	15	US-10-282-122A-43563	Sequence 43563
25	163.5	12.4	241	14	US-10-156-761-13948	Sequence 13948
26	145	11.0	231	10	US-09-557-796-32	Sequence 32
27	141	10.7	104	15	US-10-424-599-183312	Sequence 183312
28	140	10.6	202	15	US-10-282-122A-59882	Sequence 59882
29	137	10.4	234	15	US-10-282-122A-74190	Sequence 74190
30	137	10.4	238	10	US-09-557-796-18	Sequence 18
31	136.5	10.3	220	15	US-10-282-122A-49784	Sequence 49784
32	133.5	10.1	234	15	US-10-282-122A-74354	Sequence 74354
33	132	10.0	228	15	US-10-282-122A-75285	Sequence 75285
34	131	9.9	181	15	US-10-282-122A-46808	Sequence 46808
35	131	9.9	228	15	US-10-282-122A-76130	Sequence 76130
36	129.5	9.8	231	10	US-09-557-796-33	Sequence 33
37	129	9.8	228	15	US-10-282-122A-56798	Sequence 56798
38	128.5	9.7	231	15	US-10-282-122A-77900	Sequence 77900
39	127.5	9.6	232	15	US-10-282-122A-57636	Sequence 57636
40	127	9.6	230	15	US-10-282-122A-77598	Sequence 77598
41	125.5	9.5	236	15	US-10-282-122A-57141	Sequence 57141
42	122.5	9.3	228	15	US-10-282-122A-53018	Sequence 53018
43	118	8.9	213	15	US-10-282-122A-50759	Sequence 50759
44	114.5	8.7	212	15	US-10-282-122A-48161	Sequence 48161
45	112.5	8.5	233	15	US-10-282-122A-51793	Sequence 51793

ALIGNMENTS

RESULT 1

US-10-629-329A-2
; Sequence 2, Application US/10629329A
; Publication No. US2004008648A1
; GENERAL INFORMATION:
; APPLICANT: DARNAY, BRYANT G.
; TITLE OF INVENTION: METHODS AND COMPOSITIONS USING POLYNUCLEOTIDES AND
; FILE REFERENCE: POLYPTIDES OF RANK-ASSOCIATED INHIBITOR (RAIN)
; CURRENT FILING DATE: 2003-07-29
; PRIOR FILING DATE: 2002-07-29
; NUMBER OF SEQ ID NOS: 23
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 2
; LENGTH: 242
; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-629-329A-2

Query Match	100.0%	Score 1322	DB 15	Length 242
Best Local Similarity	100.0%	Pred No. 4,6e-129		
Matches 242	Conservative 0	Mismatches 0	Indels 0	Gaps 0
Qy	1	MSGCDAGEGDCCRRCAQDKHEHPYLIPELCKQPYHGLGWVTGTGGISLKHGDEIYIAP	60	
Db	1	MSGCDAGEGDCCRRCAQDKHEHPYLIPELCKQPYHGLGWVTGTGGISLKHGDEIYIAP	60	
Qy	61	SGVQKERTQPEDMFVCDINEKDISGSPSKLKKSQCTPLFNAYTMRGAGVIHTHKA	120	
Db	61	SGVQKERTQPEDMFVCDINEKDISGSPSKLKKSQCTPLFNAYTMRGAGVIHTHKA	120	
Qy	121	AVMATLLPFGREFKITHQEMIKGIKCTSGGYRVDMLVPIENTPEKGLKDRMAHA	180	
Db	121	AVMATLLPFGREFKITHQEMIKGIKCTSGGYRVDMLVPIENTPEKGLKDRMAHA	180	

QY 181 MNEYDSCAVLVRHGGVYVWGGETWEKAKTMCECYDYLFDIAVSMKKVGLDPSQLPVGENG 240
Db |||||
QY 181 MNEYDSCAVLVRHGGVYVWGGETWEKAKTMCECYDYLFDIAVSMKKVGLDPSQLPVGENG 240
Db |||||
QY 241 IV 242
Db |||||
QY 241 IV 242

RESULT 2
US-10-220-381-2
; Sequence 2, Application US/10220381
; Publication No. US20030207430A1
; GENERAL INFORMATION:
; APPLICANT: INCYTE GENOMICS, INC.
; APPLICANT: TANG, Y. Tom
; APPLICANT: LU, Dyrung Aina M.
; APPLICANT: BANDMAN, Olga
; APPLICANT: YUE, Henry
; APPLICANT: AZIMZAI, Yalda
; APPLICANT: LAL, Preeti
; APPLICANT: BURFORD, Neil
; APPLICANT: BAUGHN, Mariah R.
; TITLE OF INVENTION: HUMAN ENZYME MOLECULES
; FILE REFERENCE: PF-0763 PCT
; CURRENT APPLICATION NUMBER: US/10/220,381
; CURRENT FILING DATE: 2001-03-01
; NUMBER OF SEQ ID NOS: 52
; SOFTWARE: PERL Program
; SEQ ID NO 2
; LENGTH: 242
; TYPE: PRT
; ORGANISM: Homo sapiens
; NAME/KEY: misc feature
; FEATURE:
; OTHER INFORMATION: Incyte ID No. US20030207430A1 2116390CD1
US-10-220-381-2

Query Match 98.9%; Score 1307; DB 14; Length 242;
Best Local Similarity 99.2%; Pred. No. 1.7e-127;
Matches 240; Conservative 0; Mismatches 2; Indels 0; Gaps 0;
QY 1 MSGCDAGEGCCSRRCGAQDKHEPRYLIPELCKQFYHLGWVTGTGGGISLKHGDEIYIAP 60
Db |||||
QY 1 MSGCDAGEGCCSRRCGAQDKHEPRYLIPELCKQFYHLGWVTGTGGGISLKHGDEIYIAP 60
Db |||||
QY 61 SGVQKERIQPEDMFVCDINEKDISGSPSKLKKSOCTPLFMNAYTMRGAGAVIHTHSA 120
Db |||||
QY 121 AVMATLLFPGRFETKTHQEMIKGIKCTSGGYRYDDMLVPIIENTPEEKGLKDRMAHA 180
Db |||||
QY 121 AVMATLLFPGRFETKTHQEMIKGIKCTSGGYRYDDMLVPIIENTPEEKGLKDRMAHA 180
Db |||||
QY 181 MNEYDSCAVLVRHGGVYVWGGETWEKAKTMCECYDYLFDIAVSMKKVGLDPSQLPVGENG 240
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QY 181 MNEYDSCAVLVRHGGVYVWGGETWEKAKTMCECYDYLFDIAVSMKKVGLDPSQLPVGENG 240
Db |||||
QY 241 IV 242
Db |||||
QY 241 IV 242

RESULT 3
US-10-381-710-4
; Sequence 4, Application US/10381710
; Publication No. US20040052789A1
; GENERAL INFORMATION:
; APPLICANT: SHA, Shiken et al.
; TITLE OF INVENTION: NOVEL PROTEINS, GENES ENCODING THEM AND METHOD OF USING THE SAME
; FILE REFERENCE: 0230-0198P
; CURRENT APPLICATION NUMBER: US/10/381,710

; CURRENT FILING DATE: 2003-09-16
; NUMBER OF SEQ ID NOS: 17
; SOFTWARE: PatentIn version 3.2
; SEQ ID NO 4
; LENGTH: 242
; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-381-710-4
Query Match 98.0%; Score 1296; DB 15; Length 242;
Best Local Similarity 98.8%; Pred. No. 2.3e-126;
Matches 239; Conservative 0; Mismatches 3; Indels 0; Gaps 0;
QY 1 MSGCDAGEGCCSRRCGAQDKHEPRYLIPELCKQFYHLGWVTGTGGGISLKHGDEIYIAP 60
Db |||||
QY 1 MSGCDAGEGCCSRRCGAQDKHEPRYLIPELCKQFYHLGWVTGTGGGISLKHGDEIYIAP 60
Db |||||
QY 61 SGVQKERIQPEDMFVCDINEKDISGSPSKLKKSOCTPLFMNAYTMRGAGAVIHTHSA 120
Db |||||
QY 61 SGVQKERIQPEDMFVCDINEKDISGSPSKLKKSOCTPLFMNAYTMRGAGAVIHTHSA 120
Db |||||
QY 121 AVMATLLFPGRFETKTHQEMIKGIKCTSGGYRYDDMLVPIIENTPEEKGLKDRMAHA 180
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QY 121 AVMATLLFPGRFETKTHQEMIKGIKCTSGGYRYDDMLVPIIENTPEEKGLKDRMAHA 180
Db |||||
QY 181 MNEYDSCAVLVRHGGVYVWGGETWEKAKTMCECYDYLFDIAVSMKKVGLDPSQLPVGENG 240
Db |||||
QY 181 MNEYDSCAVLVRHGGVYVWGGETWEKAKTMCECYDYLFDIAVSMKKVGLDPSQLPVGENG 240
Db |||||
QY 241 IV 242
Db |||||
QY 241 IV 242

RESULT 4
US-10-381-710-2
; Sequence 2, Application US/10381710
; Publication No. US20040052789A1
; GENERAL INFORMATION:
; APPLICANT: SHA, Shiken et al.
; TITLE OF INVENTION: NOVEL PROTEINS, GENES ENCODING THEM AND METHOD OF USING THE SAME
; FILE REFERENCE: 0230-0198P
; CURRENT APPLICATION NUMBER: US/10/381,710
; CURRENT FILING DATE: 2003-09-16
; NUMBER OF SEQ ID NOS: 17
; SOFTWARE: PatentIn version 3.2
; SEQ ID NO 2
; LENGTH: 241
; TYPE: PRT
; ORGANISM: Mouse macrophage cell RAW 264.7
US-10-381-710-2
Query Match 93.8%; Score 1239.5; DB 15; Length 241;
Best Local Similarity 93.8%; Pred. No. 1.8e-120;
Matches 227; Conservative 9; Mismatches 5; Indels 1; Gaps 1;
QY 1 MSGCDAGEGCCSRRCGAQDKHEPRYLIPELCKQFYHLGWVTGTGGGISLKHGDEIYIAP 60
Db |||||
QY 1 MSGCQA-QGDCCSRFCGAQDKHEPRYLIPELCKQFYHLGWVTGTGGGISLKHGDEIYIAP 59
Db |||||
QY 61 SGVQKERIQPEDMFVCDINEKDISGSPSKLKKSOCTPLFMNAYTMRGAGAVIHTHSA 120
Db |||||
QY 60 SGVQKERIQPEDMFVCDINEQDISGPPASKLKKSOCTPLFMNAYTMRGAGAVIHTHSA 119
Db |||||
QY 121 AVMATLLFPGRFETKTHQEMIKGIKCTSGGYRYDDMLVPIIENTPEEKGLKDRMAHA 180
Db |||||
QY 120 AVMATLLFPGRFETKTHQEMIKGIKCTSGGYRYDDMLVPIIENTPEEKGLKDRMAHA 179
Db |||||
QY 181 MNEYDSCAVLVRHGGVYVWGGETWEKAKTMCECYDYLFDIAVSMKKVGLDPSQLPVGENG 240
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QY 180 MNEYDSCAVLVRHGGVYVWGGETWEKAKTMCECYDYLFDIAVSMKKVGLDPSQLPVGENG 239
Db |||||
QY 241 IV 242

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Db      240 IV 241
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Query Match      93.8%; Score 1239.5; DB 15; Length 241;
Best Local Similarity 93.8%; Pred. No. 1.8e-120;
Matches 227; Conservative 9; Mismatches 5; Indels 1; Gaps 1;

QY      1 MSGCDAGEGDCCRRCGAODKEHPYLIPELCKQFYHLGWTGTGGISLKHGDEIYIAP 60
Db      1 MSGCQA-QDCCSRPCGAQDKEHPRLIPELCKQFYHLGWTGTGGISLKHGNEIYIAP 59
QY      61 SGVQKERIQPEDMFVCDINEKDISGSPSKKLKKSQCTPLFMNAYTMRGAGAVIHTHSA 120
Db      60 SGVQKERIQPEDMFVCDINEQDISGPPASKKLKKSQCTPLFMNAYTMRGAGAVIHTHSA 119
QY      121 AVMATLLPGRFKEKTHQEMIKGKCTSGGYRYDDMLVPIENTPEEKLKORMAHA 180
Db      120 AVMATLLPFGQBFKEKTHQEMIKGKCTSGGYRYDDMLVPIENTPEEKLKORMAHA 179
QY      181 MNEYPDSCAVLVRHGVYVWGETWEKATMCBCYDYLFDIAVSMKKVGLDPSQLPVGENG 240
Db      180 MNEYPDSCAVLVRHGVYVWGETWEKATMCBCYDYLFDIAVSMKKVGLDPSQLPVGENG 239
QY      241 IV 242
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Db      240 IV 241

RESULT 6
US-10-437-963-195546
; Sequence 195546, Application US/10437963
; Publication No. US20040123343A1
; GENERAL INFORMATION:
; APPLICANT: La Rosa, Thomas J.
; APPLICANT: Kovalic, David K.
; APPLICANT: Zhou, Yihua
; APPLICANT: Cao, Yongwei
; APPLICANT: Wu, Wei
; APPLICANT: Boukharov, Andrey A.
; APPLICANT: Barbazuk, Brad
; APPLICANT: Li, Ping
; TITLE OF INVENTION: Rice Nucleic Acid Molecules and Other Molecules Associated with
; FILE REFERENCE: 38-21(53221)B
; CURRENT APPLICATION NUMBER: US/10/437,963
; CURRENT FILING DATE: 2003-05-14
; NUMBER OF SEQ ID NOS: 204966
; SEQ ID NO 195546
; LENGTH: 529
; TYPE: PRT
; ORGANISM: Oryza sativa

Query Match      42.6%; Score 563; DB 15; Length 256;
Best Local Similarity 51.3%; Pred. No. 6.4e-50;
Matches 117; Conservative 31; Mismatches 58; Indels 22; Gaps 7;

QY      25 RYLPELCKQFYHLGWTGTGGISLKHGDE-----IYIAPSGVQKERIQPEDMFVC 76
Db      26 RILISELCRHFYSLGWVSGTGGITIKVHDDSIKPHQILMSPSGVQKERPEPDMYVL 85
QY      77 DINEKDISGSPS-----KKLKSQCTPLFMNAYTMRGAGAVIHTHSAVMTLLFP-GR 131
Db      86 SHTGSVLSPSPKPYHPKPKCSDGCPFLMKAYEMCDAGAVIHTHSGIESCLVTMLNPLAK 145
QY      132 EFKITHQEMIKGKCTSGGYRYDDMLVPIENTPEEKLKORMAHAMNVPDSCAVL 191
Db      146 EFKITHQEMIKGK-----GHGY---DELVIPIENTAYELETESLAKIAYPKTTAVL 199
QY      192 VRRHGVYVWGETWEKATMCBCYDYLFDIAVSMKKVGLD---PSQLPV 236
Db      200 VRNHGIYVWSDSWISAKTQAECHYHFLDAAIKLHQLGLDWSSTPNHVP 247
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; FEATURE:
; OTHER INFORMATION: Clone ID: PAT_MRT4530_91484C.1.pap
US-10-437-963-195546

Query Match      43.4%; Score 573.5; DB 16; Length 529;
Best Local Similarity 46.8%; Pred. No. 1.4e-50;
Matches 123; Conservative 33; Mismatches 68; Indels 39; Gaps 7;

QY      1 MSGCDAGEGDCCRRCGAODKEHPY-----LIPELCKQFYHLGWTGTGGIS 49
Db      1 MACCGGGRGE-----GAAATESEAYLEGEAVREARLVAELCRHFYGGQWVTGTGGISIT 54
QY      50 LKHGDE-----IYIAPSGVQKERIQPEDMFVCDINEK-----DISGSPSKKLKKSOC 97
Db      55 VKANDPALPLADQLIVMSPSGVQKERVMVAEDKVLVSADGKVLSSPSVKRPWPNPKCTOC 114
QY      98 TPLFMNAYTMRGAGAVIHTHSAVMTLLPFG-REFKITHQEMIKGKCTSGGYRYD 156
Db      115 APLFMKAYLMRGAGAVIHTHSGMETCTIATMLDPGAKFRMTHMEMIKGK-----HGYR 168
QY      157 DMLVVPPIENTPEEKLKORMAHAMNVPDSCAVLVRHGVYVWGETWEKATMCBCYD 216
Db      169 DELVVPPIENTPYEYELTDSLAEALAAYPKATAVLVRNHGIYVWSDSWINAKTQAECHY 228
QY      217 LFDIAVSMKKVGLD---PSQLPV 236
Db      229 LFDAAIKLYQLGIDMTTPEHGPI 251
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; FEATURE:
; OTHER INFORMATION: Clone ID: PAT_MRT3847_112946C.1.pap
US-10-424-599-157170

RESULT 7
US-10-424-599-157170
; Sequence 157170, Application US/10424599
; Publication No. US20040031072A1
; GENERAL INFORMATION:
; APPLICANT: La Rosa, Thomas J.
; APPLICANT: Kovalic, David K.
; APPLICANT: Zhou, Yihua
; APPLICANT: Cao, Yongwei
; TITLE OF INVENTION: Soy Nucleic Acid Molecules and Other Molecules Associated with
; FILE REFERENCE: 38-21(53223)B
; CURRENT APPLICATION NUMBER: US/10/424,599
; CURRENT FILING DATE: 2003-04-28
; NUMBER OF SEQ ID NOS: 285684
; SEQ ID NO 157170
; LENGTH: 256
; TYPE: PRT
; ORGANISM: Glycine max
; FEATURE:
; OTHER INFORMATION: Clone ID: PAT_MRT3847_112946C.1.pap
US-10-424-599-157170

Query Match      42.6%; Score 563; DB 15; Length 256;
Best Local Similarity 51.3%; Pred. No. 6.4e-50;
Matches 117; Conservative 31; Mismatches 58; Indels 22; Gaps 7;

QY      25 RYLPELCKQFYHLGWTGTGGISLKHGDE-----IYIAPSGVQKERIQPEDMFVC 76
Db      26 RILISELCRHFYSLGWVSGTGGITIKVHDDSIKPHQILMSPSGVQKERPEPDMYVL 85
QY      77 DINEKDISGSPS-----KKLKSQCTPLFMNAYTMRGAGAVIHTHSAVMTLLFP-GR 131
Db      86 SHTGSVLSPSPKPYHPKPKCSDGCPFLMKAYEMCDAGAVIHTHSGIESCLVTMLNPLAK 145
QY      132 EFKITHQEMIKGKCTSGGYRYDDMLVPIENTPEEKLKORMAHAMNVPDSCAVL 191
Db      146 EFKITHQEMIKGK-----GHGY---DELVIPIENTAYELETESLAKIAYPKTTAVL 199
QY      192 VRRHGVYVWGETWEKATMCBCYDYLFDIAVSMKKVGLD---PSQLPV 236
Db      200 VRNHGIYVWSDSWISAKTQAECHYHFLDAAIKLHQLGLDWSSTPNHVP 247
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; FEATURE:
; OTHER INFORMATION: Clone ID: PAT_MRT4530_91484C.1.pap
US-10-437-963-195546
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RESULT 11
US-10-767-701-45914

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RESULT 9
US-10-425-114-64486
; Sequence 64486, Application US/10425114
; Publication NO. US20040034888A1
; GENERAL INFORMATION:
; APPLICANT: Liu, Jingdong
; APPLICANT: Zhou, Yihua
; APPLICANT: Kovalic, David K.
; APPLICANT: Screen, Steven E.
; APPLICANT: Tabaska, Jack E
; APPLICANT: Cao, Yongwei
; TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With
; TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
; FILE REFERENCE: 38-21(53313)B
; CURRENT APPLICATION NUMBER: US/10/425,114
; CURRENT FILING DATE: 2003-04-28
; NUMBER OF SEQ ID NOS: 73128
; SEQ ID NO 64486
; LENGTH: 524
; TYPE: PRT
; ORGANISM: Zea mays
; FEATURE:
; OTHER INFORMATION: Clone ID: LIB3060-104-F8_FLI.pep
US-10-425-114-64486

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; Sequence 45914, Application US/10767701
; Publication No. US20040172684A1
; GENERAL INFORMATION:
; APPLICANT: Kovalic, David K.
; APPLICANT: Zhou, Yihua
; APPLICANT: Cao, Yongwei
; TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With
; FILE REFERENCE: 38-21(53535)B
; CURRENT APPLICATION NUMBER: US/10/767,701
; CURRENT FILING DATE: 2004-01-29
; NUMBER OF SEQ ID NOS: 63128
; SEQ ID NO 45914
; LENGTH: 517
; TYPE: PRT
; ORGANISM: Sorghum bicolor
; FEATURE:
; OTHER INFORMATION: Clone ID: SORBI-28MAY03-C2601_1.pap
US-10-767-701-45914

Query Match 41.8%; Score 553; DB 16; Length 517;
Best Local Similarity 46.1%; Pred. No. 1.8e-48;
Matches 117; Conservative 34; Mismatches 71; Indels 32; Gaps 7;

Qy 9 GDC-CSRRCQAQDKB-----HPRYLIPELCKQFYHLGWVTGGGSLKHGDE--- 55
Db 4 GGCCEAAVAGTASEAYLEGEVPREARELVAELCRHFYAQGWVTGGGSIYKVNDRPVP 63
Qy 56 ----IYIAPSGVQKRIQPEDMFVCDINEKDISGPS---PSKKLKKSQCTPLFMNAYT 106
Db 64 LADRLIWNPSGVQKRVMAEDMYMAADGKVLSPVAKPKNPKCTDCAPLFMKAYL 123
Qy 107 MRGAGAVIHTHSKAAVMATLLPFG-REFKITHQEMIKGIKKCTSGGYRYDDMLVVPPIE 165
Db 124 MRGAGAVIHTSHGMECTIATMLNPGAKEFRMTHMEMIKGIG-----HGYRDELVIPIVE 177
Qy 166 NTPBEKGLKDRMAHAMNEYPDSCAVLVRHGVYVWGETWEKATMCBCYDYLFDIAVSMK 225
Db 178 NTPYEYELTSLSEIAAAYPKATAVLVRNHGIVYWGDSWINAKTQAECHYLLDACIKLY 237
Qy 226 KVGLED---PSQLPV 236
Db 238 QLGIDWTTPEHGPI 251

RESULT 12
US-10-424-599-273717
; Sequence 273717, Application US/10424599
; Publication No. US20040031072A1
; GENERAL INFORMATION:
; APPLICANT: La Rosa, Thomas J
; APPLICANT: Kovalic, David K
; APPLICANT: Zhou, Yihua
; APPLICANT: Cao, Yongwei
; TITLE OF INVENTION: Soy Nucleic Acid Molecules and Other Molecules Associated With
; FILE REFERENCE: 38-21(53223)B
; CURRENT APPLICATION NUMBER: US/10/424,599
; CURRENT FILING DATE: 2003-04-28
; NUMBER OF SEQ ID NOS: 285684
; SEQ ID NO 273717
; LENGTH: 522
; TYPE: PRT
; ORGANISM: Glycine max
; FEATURE:
; OTHER INFORMATION: Clone ID: PAT_MRT3847_89188C.1.pap
US-10-424-599-273717

Query Match 41.5%; Score 549; DB 15; Length 522;
Best Local Similarity 51.3%; Pred. No. 4.9e-48;
Matches 117; Conservative 29; Mismatches 60; Indels 22; Gaps 7;

Qy 25 RYLIPELCKQFYHLGWVTGGGSLK-HGDE-----IYIAPSGVQKRIQPEDMFVC 76
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Db 29 RALMAELCRHFYTLGWVTGGGSLKSMKVHDDSI PRPQQLILMAPSGVQKERPEDEMYVL 88
Qy 77 DINEKDISGPSPS---KKLKKSQCTPLFMNAYTMRGAGAVIHTHSKAAVMATLLP-GR 131
Db 89 SHSGSVLSAPSPKWPHPKPCDCDPLFKKAYENRDAANVPHSHGIESCLVTMINPLSK 148
Qy 132 EFKITHQEMIKGIKKCTSGGYRYDDMLVVPPIENTPEEKGKDRMAHAMNEYPDSCAVL 191
Db 149 EFRITHMEMIKGIG--GHGY--DELVVPPIENTAYEYQLTESFAKAIEDYPKATAVL 202
Qy 192 VRRHGVYVWGETWEKATMCBCYDYLFDIAVSMKVGLED---PSQLPV 236
Db 203 VRNHGVFWGDSWISAKTQSECYHYLFDALKLHQMGLDWSPTPNHGPI 250

RESULT 13
US-10-425-114-46271
; Sequence 46271, Application US/10425114
; Publication No. US20040034888A1
; GENERAL INFORMATION:
; APPLICANT: Liu, Jingdong
; APPLICANT: Zhou, Yihua
; APPLICANT: Kovalic, David K.
; APPLICANT: Screen, Steven E.
; APPLICANT: Tabaska, Jack E.
; APPLICANT: Cao, Yongwei
; TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With
; FILE REFERENCE: 38-21(53313)B
; CURRENT APPLICATION NUMBER: US/10/425,114
; CURRENT FILING DATE: 2003-04-28
; NUMBER OF SEQ ID NOS: 73128
; SEQ ID NO 46271
; LENGTH: 540
; TYPE: PRT
; ORGANISM: Glycine max
; FEATURE:
; OTHER INFORMATION: Clone ID: 701207771_FLI.pap
US-10-425-114-46271

Query Match 41.5%; Score 549; DB 15; Length 540;
Best Local Similarity 51.3%; Pred. No. 5.1e-48;
Matches 117; Conservative 29; Mismatches 60; Indels 22; Gaps 7;

Qy 25 RYLIPELCKQFYHLGWVTGGGSLK-HGDE-----IYIAPSGVQKRIQPEDMFVC 76
Db 47 RALMAELCRHFYTLGWVTGGGSLKSMKVHDDSI PRPQQLILMAPSGVQKERPEDEMYVL 106
Qy 77 DINEKDISGPSPS---KKLKKSQCTPLFMNAYTMRGAGAVIHTHSKAAVMATLLP-GR 131
Db 107 SHSGSVLSAPSPKWPHPKPCDCDPLFKKAYENRDAANVPHSHGIESCLVTMINPLSK 166
Qy 132 EFKITHQEMIKGIKKCTSGGYRYDDMLVVPPIENTPEEKGKDRMAHAMNEYPDSCAVL 191
Db 167 EFRITHMEMIKGIG--GHGY--DELVVPPIENTAYEYQLTESFAKAIEDYPKATAVL 220
Qy 192 VRRHGVYVWGETWEKATMCBCYDYLFDIAVSMKVGLED---PSQLPV 236
Db 221 VRNHGVFWGDSWISAKTQSECYHYLFDALKLHQMGLDWSPTPNHGPI 268

RESULT 14
US-10-425-115-253964
; Sequence 253964, Application US/10425115
; Publication No. US20040214272A1
; GENERAL INFORMATION:
; APPLICANT: La Rosa, Thomas J.
; APPLICANT: Kovalic, David K.
; APPLICANT: Zhou, Yihua
; APPLICANT: Cao, Yongwei
; TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With
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Wed Feb 2 10:19:19 2005

QY 112 AVIHTSKAAVMATLLPPG-REFKITHOEMIKGIKCTSGGYRYDDMLVPIIENTPEE 170
Db 71 AVIHTSHGIECTIATMLIPGAKPRVTHMEMIKIGK-----HGYHDELVIPIIENTPYE 124
QY 171 KGLKDRMAHAMNEYPDSCAVLVRHGHVYVWGCTWEKAKTMCBCYDYLFDIAVSMKKVGLD 230
Db 125 YELTDSLSEALAAYPKATAVLVRNHGIYVWGESWINAKTQACFYHYLLDACIKLYQLGID 184

Search completed: February 1, 2005, 14:33:32
Job time : 149 secs

Query Match 38.9%; Score 514.5; DB 17; Length 594;
Best Local Similarity 36.0%; Pred. No. 2.3e-44;
Matches 118; Conservative 31; Mismatches 68; Indels 111; Gaps 7;
QY 2 SGCDAGEGDCSRRRCAGDKE-----HPRVLIPELCKQFYHLGWVGTGGGSLKH 52
Db 4 SGCS-----CEAAVGAASEAYLEGAPVREARELVAELCRHFYAQQGVGTGGSIYKV 57
QY 53 GDE-----IYIAPSGVOKERIOPEDMFVCDINEKDISGPS-----PSKKLKSQCTPL 100
Db 58 NDPTVPLADRLIWMSPSGVOKERWVAEDMYMAADGKVLAPVAKPWPKPKCTDCAPL 117
QY 101 FMNAYTMRGAGAVIHTHSCAAVMATLLPPG-REFKITHOEMIKGIKCTSGGYRYDDML 159
Db 118 FMKAYLMRGAGAVIHTSHGIECTIATMLIPGAKPRVTHMEMIKIGK-----HGYHDEL 171
QY 160 VPIIENTPEEKLKDRMAHAMNEYPDSCAVLVRHGHVYVWGCTWEKAKT----- 210
Db 172 VPIIENTPYEELTDSLSEALAAYPKATAVLVRNHGIYVWGESWINAKTQACFGFRDQ 231
QY 211 ----- 210
Db 232 IKDFIWMTLKPELDHASFRKNTMYLHIFMGSLILQSFVIIITDVLVAGGIWGRNS 291
QY 211 -----CECYDYLFDAIVSMKKVGLD 230
Db 292 LTPAFCREACFYHYLLDACIKLYQLGID 319

RESULT 15
US-10-425-114-61505
; Sequence 61505, Application US/10425114
; Publication No. US20040034888A1
; GENERAL INFORMATION:
; APPLICANT: Liu, Jingdong
; APPLICANT: Zhou, Yihua
; APPLICANT: Kovalic, David K.
; APPLICANT: Screen, Steven E
; APPLICANT: Tabaska, Jack E
; APPLICANT: Cao, Yongwei
; TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated with
; TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
; FILE REFERENCE: 38-21(53313)B
; CURRENT APPLICATION NUMBER: US/10/425,114
; CURRENT FILING DATE: 2003-04-28
; NUMBER OF SEQ ID NOS: 73128
; SEQ ID NO 61505
; LENGTH: 459
; TYPE: PRT
; ORGANISM: Zea mays
; FEATURE:
; OTHER INFORMATION: Clone ID: LIB143-005-F3_FLI.pep
US-10-425-114-61505

Query Match 35.4%; Score 467.5; DB 15; Length 459;
Best Local Similarity 51.7%; Pred. No. 1.3e-39;
Matches 93; Conservative 27; Mismatches 49; Indels 11; Gaps 3;
QY 56 IYIAPSGVOKERIOPEDMFVCDINEKDISGPS-----PSKKLKSQCTPLFMNAYTMRGAG 111
Db 11 IWMSPSGVOKERWVAEDMYMAADGKVLAPVAKPWPKPKCTDCAPLFMKAYLMRGAG 70